Microbial enzyme activity and carbon cycling in grassland soil aggregates

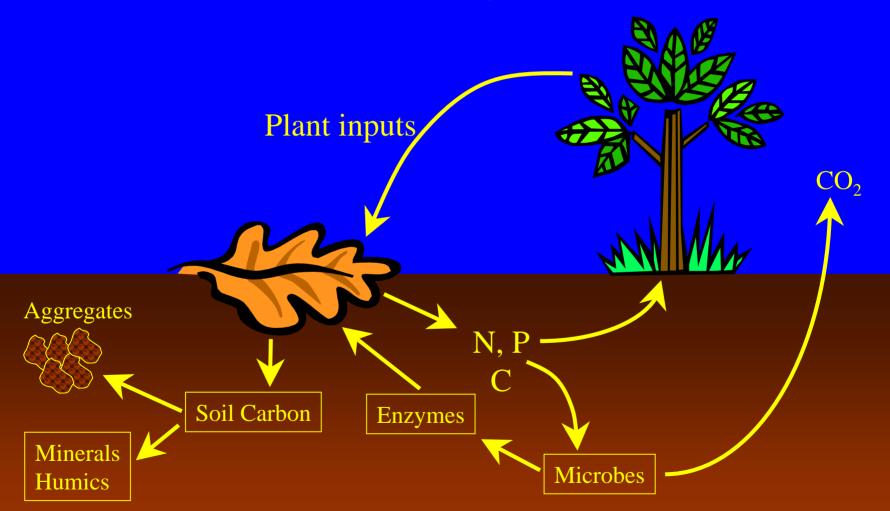
Steve Allison GCEP Workshop August 17-20, 2003

Primary research question

 What controls the rate of carbon and nutrient cycling in soil?

How do soil microorganisms make a living?

The soil system



Soil Enzymes

 Produced by microbes to degrade and assimilate complex compounds

 First step to convert soil organic matter into CO₂ and inorganic nutrients

 Can affect soil carbon sequestration and CO₂ release

Enzymes and their functions

Enzyme	Function
Acid phosphatase	Releases PO ₄ ³⁻
Urease	Releases NH ₄ ⁺ from urea
Chitinase	Releases chitin monomers
Beta-glucosidase	Degrades cellulose
Cellobiohydrolase	Degrades cellulose
Polyphenol oxidase	Degrades lignin

An enzymic 'latch' on a global carbon store

A shortage of oxygen locks up carbon in peatlands by restraining a single enzyme.

istorically, northern peatlands have removed carbon dioxide from the atmosphere faster than it has been rereleased, so they now contain 20–30% of the world's soil carbon stock¹ (the equivalent of over 60% of the atmospheric carbon pool²). Here we show that the anaerobic conditions in peatlands prevent the enzyme phenol oxidase from eliminating phenolic compounds that inhibit biodegradation. This indicates that oxygen limitation on a single peatland enzyme may be all that prevents the rerelease of a major store of global carbon into the atmosphere, with potentially serious implications for future global warming.

Mechanisms proposed to account for the slow decomposition rates in peatlands include the effects on microbial metabolism of low oxygen availability, low pH, low nutrient supply and low temperatures. But decomposition can be highly efficient in

Table 1 Effects on enzyme activities		
	Control	Manipulated
Effect of oxygen		
Sulphatase	66±2.3	35±1.4
Phosphatase	571 ± 2.4	387±7.9
β-Glucosidase	237 ± 2.3	177±12
Phenol oxidase	615 ± 93	4,350±27
Effect of increasi	ng phenol oxida	se abundance
Phenolics (μg I ⁻¹)	1,985 ± 55.4	1,444±9.9
β-Glucosidase	1,677±280	10,111±380
Effect of phenolic	removal on hy	drolase activity
Sulphatase	579 ± 36	849±43
Phosphatase	3,707 ± 25	4,369±180
β-Glucosidase	1,723 ± 120	2,183 ± 180
Xylosidase	116±2.5	134±5
Chitinase	243 ± 14	296±3.5

Phenol oxidase activity (nmol 2-carboxy-2,3-dihydroindole-5,6-quinone formation min⁻¹ per g peat), hydrolase activities (nmol methylumbelliferone formation min⁻¹ per g peat) and phenolic compound concentrations (μ g l⁻¹) are reported as mean \pm s.e.

(Table 1). Lower water-tables, which are

field survey (r=0.61, P<0.05) that every doubling in phenol oxidase activity was accompanied by an approximate doubling in CO₂ production.

Taken together, our findings support the idea that oxygen constraints on a single enzyme, phenol oxidase, can minimize the activity of hydrolytic enzymes responsible for peat decomposition. This has profound implications in the context of climate change as a feedback to the process of intensified carbon loss. Increased peat aeration, as a result of droughts predicted by certain climate-change models¹³, has the potential to eliminate a critical mechanism restricting the re-release of CO₂ to the atmosphere. As such, phenol oxidase could be considered to represent a fragile 'latch' mechanism holding in place a vast carbon store of 455 gigatonnes.

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Questions for ANL work

What controls soil carbon storage?

 Why is so much carbon stored within soil aggregates?

 Are enzymes (or lack of enzymes) responsible?

CONCEPTUAL DIAGRAM OF AGGREGATE HIERARCHY

From Jastrow and Miller, 1998, In Soil Processes and the Carbon Cycle, CRC Press.



Microaggregates

- Dead fungi
- **O** Clay with microbes
- Clay particles



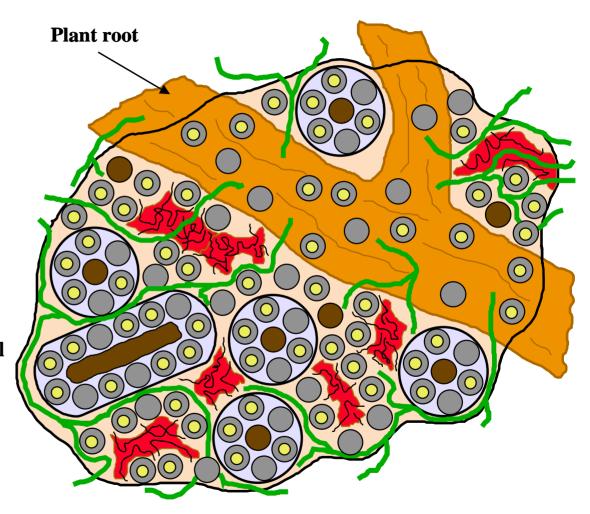
Decaying plant material



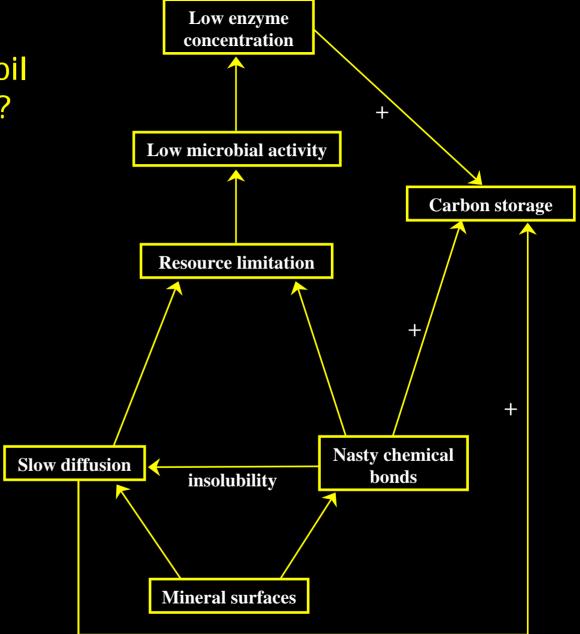
Symbiotic fungi

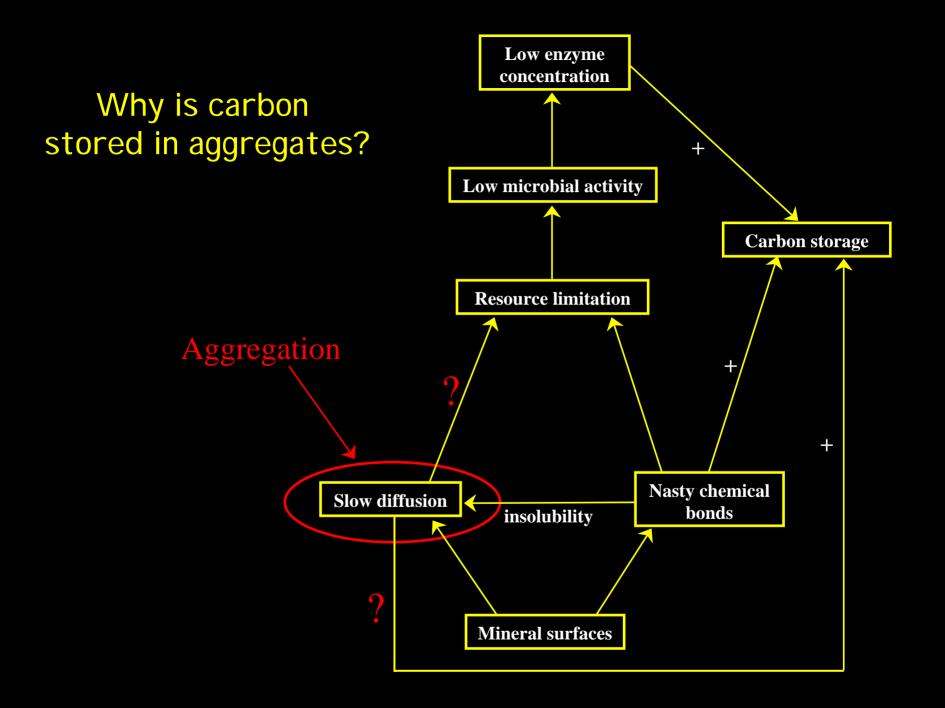


Binding agents: "glue"



What controls soil carbon storage?



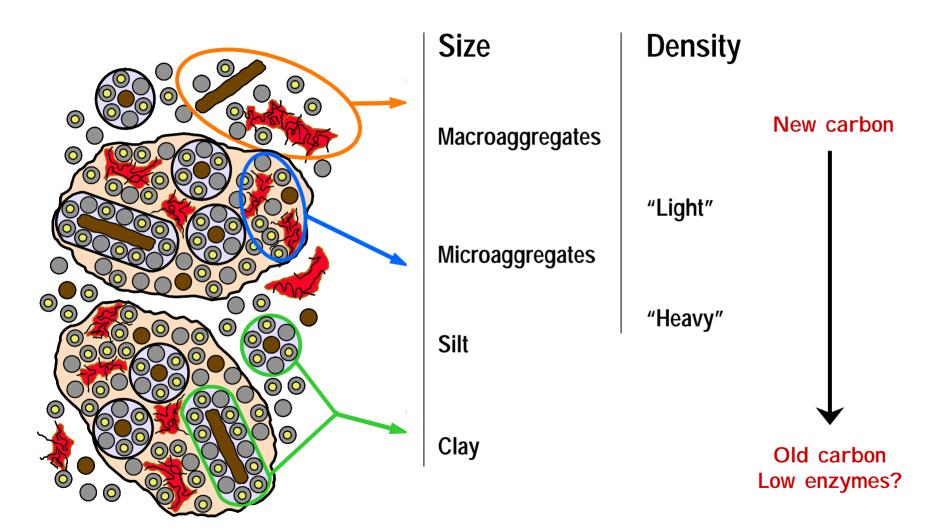


Experimental Design

Separation Enzyme activities Planted pasture Composite Agricultural Restored prairie Native prairie

Increasing soil C and aggregation

Fractionation of Soil Organic Matter Based on Aggregate Hierarchy



J.D. Jastrow, Argonne National Laboratory

Predictions

Fraction	Prediction
Light organic matter	Accessible substrates, high microbial activity, high enzyme activity
Macroaggregates	Includes some light organic matter; above average enzyme activity
Microaggregates	Physically shielded, below-average enzyme activity
Silt	Low enzyme activity
Clay	Lowest enzyme activity

No actual results yet BUT...

I can use models to make them up!



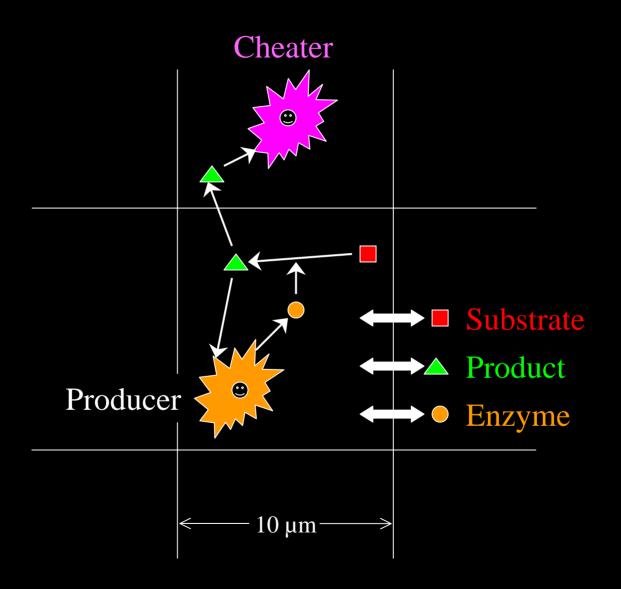
Enzyme production model

- Spatially explicit
- Driven by microbes
- Microbes produce enzymes to get carbon and nutrients

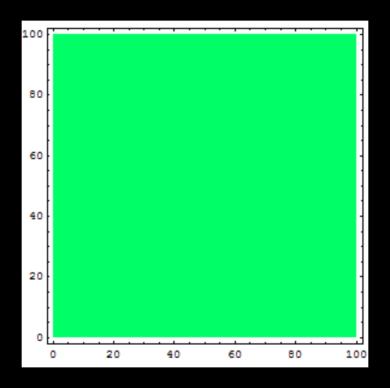
Outputs:

- Population sizes
- Enzyme, substrate, and product concentrations
- CO₂ production rates

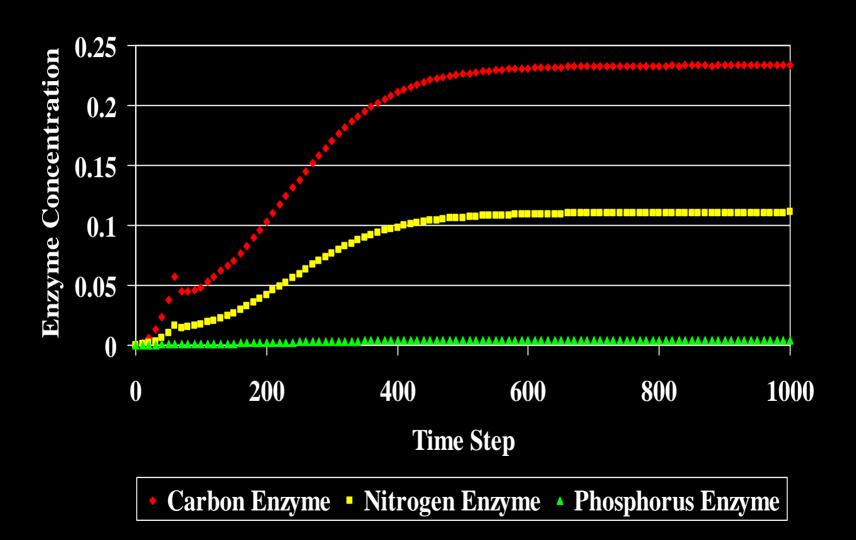
Grid-based enzyme production model



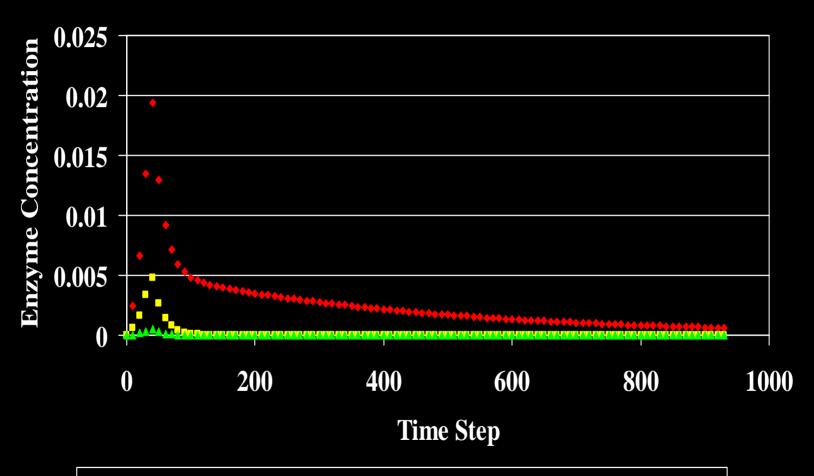
Example: Product concentrations over time on the grid



Enzyme concentrations: outside aggregates



Enzyme concentrations in aggregates



Carbon Enzyme
Nitrogen Enzyme
Phosphorus Enzyme

Concluding thoughts

- Microbes control rates of decomposition and nutrient release
- They need enzymes to do this
- Enzyme activities should reflect the soil conditions within aggregates
- Old soil carbon should not be associated with high enzyme activity

Thanks:

- DOE-GCEP
- Julie Jastrow
- Peter Vitousek